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CELLULASE PRODUCTION BY MICROORGANISMS ISOLATED FROM LAGUNA BLANCA, POTOSÍ-BOLIVIA

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ABSTRACT

Bacterial strains isolated from water samples collected from Laguna Blanca, Potosi-Bolivia and deposited at Biotechnology Center, Cochabamba-Bolivia, were subjected to cellulase enzyme production studies. Among 71 strains, 32 were selected as positive in liquid medium using filter paper as sole carbon source. Nine strains reported higher cellulase activity and the effect of NaCl concentration and pH on microorganism growth was evaluated. Thus, 7 strains were identified as halophiles meanwhile 2 were halotolerant. According to the medium pH value, 5 strains were neutrophiles and 4 alkalotolerants. Strains coded as LB-4 and LB-8 were selected as the best enzyme producers. Furthermore, cellulase production using forest products and agro-industrial residues were positive for different types of paper and pine wood. Nevertheless, higher activity values were detected using banana peel and sugarcane bagasse. The strain LB-8 was selected as the best cellulase producer under the conditions used in this study with 2.774 U/ml enzyme activity. Phylogenetic analyses showed 99.99 % sequence similarity between strain LB-8 and *Bacillus pumilus*.

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RESUMEN

Las cepas bacterianas aisladas de muestras de agua recolectadas de Laguna Blanca, Potosí-Bolivia, y depositadas en el Centro de Biotecnología, Cochabamba-Bolivia, fueron sometidas a estudios de producción de celulasas. Entre 71



cepas, 32 se seleccionaron como positivas en medio líquido utilizando papel de filtro como única fuente de carbono. Nueve cepas reportaron mayor actividad de celulasas y se evaluó el efecto de la concentración de NaCl y el pH sobre el crecimiento de microorganismos. Siete cepas se identificaron como halófilas mientras que 2 fueron halotolerantes. Según el valor de pH del medio de cultivo, 5 cepas fueron neutrófilas y 4 alcalotolerantes. Las cepas codificadas como LB-4 y LB-8 fueron seleccionadas como mejores productoras de celulasas. Además, la producción de celulasas utilizando productos forestales y residuos agroindustriales fue positiva para diferentes tipos de papel y madera de pino, sin embargo, se detectaron valores de actividad más altos con cáscara de plátano y bagazo de caña de azúcar. Se seleccionó la cepa LB-8 como la mejor productora de celulasas bajo las condiciones utilizadas en este estudio con una actividad enzimática de 2.774 U/ml. Los análisis filogenéticos mostraron una similitud de secuencia del 99,99 % entre la cepa LB-8 y *Bacillus pumilus*.

INTRODUCTION

Cellulose is an unbranched glucose polymer, composed of anhydro-D-glucose units linked by 1,4- β -D-glucoside bonds, which can be hydrolyzed by cellulolytic enzymes produced by both bacteria and fungi [1]. Cellulose can be considered as the most abundant and biologically renewable resource for bioconversion processes. Thus, the progress in biotechnology of cellulases and related enzymes is a promising way for obtaining sugars from lignocellulosic materials. The possibility of enzymatically converting the cellulose present in forest products and agro-industrial waste into useful products like bioethanol is emerging as an alternative to fossil fuels. The necessity of cost reduction in the bioethanol production industry led to renewed interest in these types of enzymes [2-4].

The microbial biodiversity of extreme environments represents a promising source for more stable, highly active, and specific enzymes. Lakes and lagoons distributed in the Andes, are subjected to extreme temperature variations, high UV radiation levels as well as a wide range of pH values and metal concentrations. In order to survive microorganisms isolated from these environments have developed strong metabolic systems for adaptation. Laguna Blanca is located in Bolivian highlands at 4,300 m above sea level, in Potosí-Bolivia, into a xeric volcanic landscape of the High Andean Ecoregion, its coloration is quite white caused by the high content of salt and minerals in its waters [5].

Enzymes produced by extremophiles are currently under study due to their properties and potential applications. Bacterial cellulases have been reported over the past decades; nonetheless, new strains are still under studies since new microorganisms represent a promising source of new enzymes which may be more stable under industrial process conditions [6, 7]. The present study evaluates the bacterial biodiversity of Laguna Blanca, Potosi-Bolivia, for cellulase production using low price substrates.

EXPERIMENTAL

Chemicals

Agro-industrial waste was provided by local feed plants: wheat straw, corn stalk, corn leaves, and rice husks. Meanwhile, banana peel and sugarcane waste were from local markets. All other chemicals were purchased either from Merck or Sigma. All other chemicals used in the present study were of analytical grade.

Isolation and selection of cellulase-producing microorganisms

Water samples were collected from Laguna Blanca shoreline in 100 ml sterile flasks from the surface water body (22°46' S, 67°47' W), Potosí-Bolivia, and stored at 4 °C before processing. The samples were enriched on growth liquid medium for chemotrophic microorganisms (QM) containing (% w/v): [Yeast extract - 2, peptone - 1, $\text{MgCl}_2 \cdot 6\text{H}_2\text{O}$ - 0.875, $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ - 0.6, KCl - 0.125, $\text{C}_6\text{H}_5\text{Na}_3\text{O}_7$ - 0.1, $\text{CaCl}_2 \cdot 2\text{H}_2\text{O}$ - 0.025, NaBr - 0.0125%, NaHCO_3 - 0.005, NaCl 0 to 15] and for heterotrophic (HM) microorganisms containing (% w/v): [Yeast extract - 1, peptone - 0.5, glucose - 0.1, $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ - 0.025, KCl - 0.05, $\text{CaCl}_2 \cdot 2\text{H}_2\text{O}$ - 0.009, NaBr - 0.006, NaCl - 0 to 15] the pH was adjusted to 7 using a 2M NaOH solution. Cellulase-producing microorganisms were selected in liquid medium (QM/HM) supplemented with filter paper, 1 % (w/v), as sole carbon source in a rotary incubator shaker (Innova 4430, New Brunswick Scientific) at 200 rpm orbital agitation speed and 30 °C. Cell growth was monitored by measuring the optical density in a spectrophotometer (UV-Vis 1601, Shimadzu) at 600 nm, and stock cultures were maintained on isolation medium at 4 °C and as glycerol stock at -20 °C.

Effect of NaCl concentration and pH on microorganism growth



Solid culture medium (QM/HM), with added agar 1,5% (w/v) and pH 7, was prepared at different NaCl % (w/v) concentrations: 0, 5, 10 and 15 and inoculated with the best cellulolytic enzyme producer. The pH effect on the microorganism growth was evaluated with the best NaCl % (w/v) concentration previously determined, at pH values of 6, 7, 7.5, 8 and 9. The cultures were grown at 30 °C for 12 hours and positive or negative bacterial growth was observed in solid medium.

Cellulase production using forest products and agro-industrial waste

Enzyme production was studied in liquid culture medium (QM/HM) supplemented with filter paper 1% (w/v) as sole carbon source and incubated at 30 °C and 200 rpm orbital agitation speed for 48 hours. At 12 hours intervals, aliquots were withdrawn and centrifuged at $7.155 \times g$ for 10 min. The cell free supernatant was analyzed for cellulolytic activity.

Those strains selected as best cellulase producers were cultivated in liquid culture medium (QM/HM) supplemented with 1% (w/v) organic waste individually: forest products like pine wood and different types of paper (newspaper, printing paper, office paper and filter paper) as well as agro-industrial waste like wheat straw, corn stalk, corn leaves, banana peel, sugarcane, and rice husks as carbon source. The cultures were grown at 30 °C, 200 rpm for 15 hours, samples were taken each 3 hours and the cell-free supernatant after centrifugation was analyzed for cellulase enzyme activity, measurements were performed in duplicate. All the substrates used were rinsed twice with distilled water, dried at 50°C, stored at -20°C until use and sterilized with the medium.

The optimum pH of the enzyme crude extract of the selected strain was measured at 37°C by using buffer solutions of different pH values. The following buffers (50mM) were used: potassium phosphate (pH 6.0-8.0), Tris-HCl (pH 8.0-9.0) and glycine/NaOH (pH 9.0-12). In order to determine the optimum temperature for the enzyme extract, cellulase activity was assayed at 20, 30, 37, 40, 50 and 60 °C.

Statistical analysis

Mann-Whitney U Test (non-parametric data) was used for statistical analyses for all the organic waste substrates tested. A Wilcoxon Signed Ranks Test was used to evaluate the groups of organic waste in this study [8].

Microorganisms identification

In order to identify the isolated species, deoxyribonucleic acid (DNA) extraction, 16S rDNA gene amplification, and sequencing of the PCR products were carried out at Bio Basic Inc. (Canada). The analysis of the DNA sequences was performed with the BLAST server in the National Center for Biotechnology Information (<http://www.ncbi.nlm.nih.gov>) using the BLAST algorithm and the BLASTN software for the comparison of a nucleotide query sequence against a nucleotide sequence database. 16S rDNA sequence analysis was performed with the MEGA X software package using the neighbor-joining method [9]. For the phylogenetic trees, only sequences from the type strains whose names have been validly published were taken into account.

Analytical methods

Cellulase assay

The cellulase enzyme activity was determined by measuring the release of reducing sugars from filter paper (1% w/v) by dinitro salicylic acid (DNS) method. A 100 µl of the enzyme extract was mixed with 200 µl of DNS and incubated in a boiling water bath for 10 min, the reaction was stopped in a cold water bath and then the absorbance at 540 nm was measured on the spectrophotometer. A blank with 100 µl of distilled water instead of the enzyme extract was used in all the different tests [10]. One unit of cellulase activity was defined as amount of enzyme required to release 1µmol of glucose from filter paper in 1 minute under standard conditions.

Reducing sugars determination

Estimation of total reducing sugars in the enzymatic hydrolysate of biomass was quantified by DNS method [10].

RESULTS AND DISCUSSION



Cellulase-producing microorganism selection

A total of 71 bacterial strains were isolated from Laguna Blanca liquid samples and grown in liquid medium supplemented with filter paper as sole carbon source. From 32 strains which gave positive results for cellulase production, 9 reported higher cellulase activity in liquid media. The strains coded as LB-1, LB-2, LB-4, LB-6, LB-7, LB-8, LB-13, LB-17 y LB-22 were selected for further studies. Successful screening of novel cellulase-producing strains represents a strategy for multifunctional cellulases production with broader substrate utilization as well as wider temperatures and pH activity range conditions [11].

Effect of NaCl concentration and pH on microorganism growth

The 9 strains selected were grown at different NaCl concentrations ranging for 0 to 15% (w/v) from those, 7 strains: LB-4, LB-6, LB-7, LB-8, LB-13, LB-17 and LB-22 were identified as halophilic and 2 strains: LB-1 and LB-2 as halotolerant. Microorganisms that grew between 3 and 15% (w/v) NaCl are considered halophiles and those that grow well without NaCl and are able to grow in the presence of NaCl are defined as halotolerant. Enzymes produced by these microorganisms have proved to present remarkable properties of commercial interest due to its stability at different salt concentrations and pH values [12]. Screening studies in saline habitats have reported numerous microorganisms as source of extremophilic enzymes with promising applications in biotechnological processes mainly because they are subjected to different environmental stresses simultaneously [13].

Cell growth classification according to different pH values was carried out considering alkaliphilic microorganisms those that grow optimally in pH values greater than 9, meanwhile neutrophiles grow optimally in pH values between 6 and 9. Alkalotolerants are those microorganisms that grow optimally near neutral pH but also tolerate pH values greater than 9 [13]. Five strains: LB-2, LB-4, LB-6, LB-7 and LB-8 are neutrophiles meanwhile 4 strains: LB-1, LB-13, LB-17 and LB-22 are alkalotolerants. Extracellular salt-tolerant enzymes of moderately halophilic bacteria and alkaliphilic/alkalotolerant enzymes from alkaliphilic bacterial strains offer broad application in biotechnological processes in food, laundry detergent, leather, and paper/pulp industries. Thus, a considerable effort has been dedicated to the study of such enzymes due to its remarkable properties [14-16].

The 9 selected strains were cultivated in liquid media supplemented with filter paper, 1% (w/v), as sole carbon source. Higher enzyme activity values were registered at 12 hours and strains LB-4 and LB-8 were selected for further studies (Fig. 1).

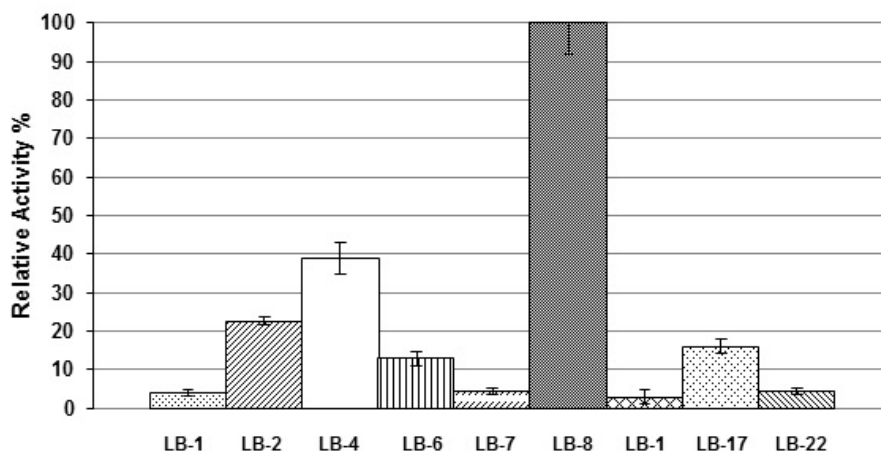


Figure 1. Cellulase relative activity (%) from strains LB-1, LB-2, LB-4, LB-6, LB-7, LB-8, LB-13, LB-17 and LB-22 using filter paper, 1% (w/v), as carbon source after 12 hours incubation. Error bars represent the mean \pm 2SD of duplicate samples.

Cellulase production using organic waste: forest products and agro-industrial waste

The 2 strains selected as best cellulase producers (LB-4 and LB-8) were cultivated in liquid culture media supplemented with 1% (w/v) organic waste: forest products and agro-industrial waste as carbon source. However, a Mann Whitney U Test indicated that the difference between cellulase activity of the selected strains on organic waste tested was non-significant ($U=48$, $p=0.206$). After 9 hours cultivation in forest products, strain LB-4 showed highest enzyme activity values using printing paper, 0.071 (U/ml), and pine wood, 0.114 (U/ml), meanwhile for strain LB-



8 the highest values were registered when using newspaper, 0.150 (U/ml), and pine wood, 0.148 (U/ml), (Fig. 2). Paper recycling represents a subsection of paper industry oriented towards re-use and sustainability. The application of cellulases for paper recycling has proved to improve defibering and ink particle detachment [17, 18]. Strains LB-4 and LB-8 were able to growth using different types of papers as well as pine wood thus they can be applied in industrial paper processes.

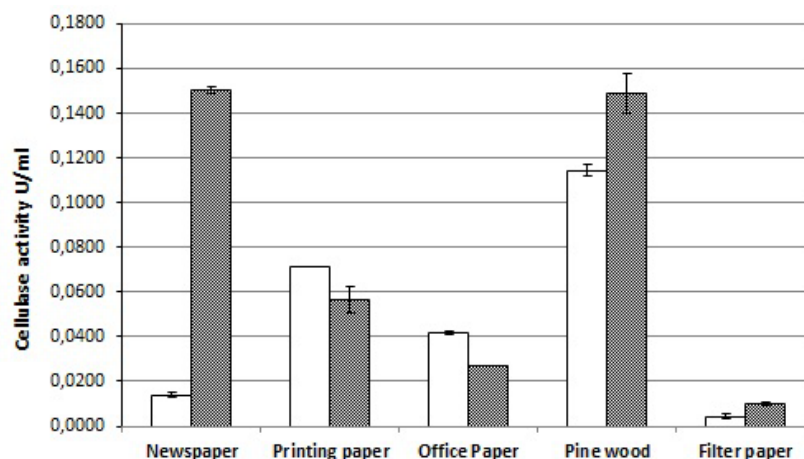


Figure 2. Cellulase activity (U/ml) for strains (□) LB-4 and (▨) LB-8 using 1% (w/v) forest products as carbon source. Error bars represent the mean \pm 2SD of duplicate samples.

In the case of agro-industrial waste, the highest enzyme activity for LB-4 and LB-8 was detected using banana peel, 0.944 (U/ml) and 2.774 (U/ml), respectively. In the same way, using sugarcane bagasse presented activity values of 1.849 and 2.774 (U/ml) for both LB-4 and LB-8 (Fig. 3). Cellulose is a linear polysaccharide polymer that greatly differs from one source to another in structure. Therefore, the hydrolysis of this material depends on the type of residue. A Wilcoxon Signed-Ranks Test indicated that there is a significant difference in the enzyme activity by the two strains ($Z = 2.55$, $p = 0.011$) when agro-industrial organic waste was tested for cellulases. Strain LB-8 registered higher enzyme activity values when compared to strain LB-4, 2.774 (U/ml). Therefore, strain LB-8 was selected for phylogenetic analysis. Cellulase production using agro-industrial waste is an economical option, since harvesting and processing agricultural feedstock produce an enormous quantity of solid lignocellulosic residue that can be considered one of the major environmental pollutants due to their unchecked accumulation. Microorganisms degrade lignocellulosic material due to their highly efficient enzymatic system, thus, several studies are directed towards the production of hydrolytic enzymes using low-cost substrates that can be applied in biotechnology and biorefinery approaches for fuel generation, chemicals production as well as energy sources [19-21].

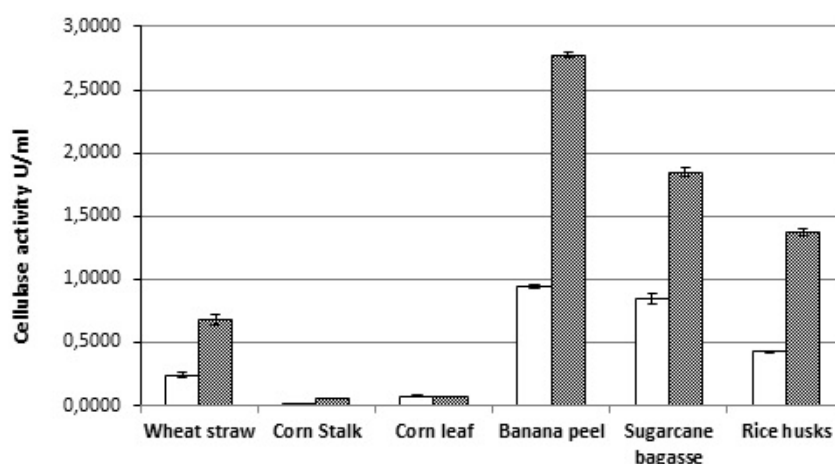


Figure 3. Cellulase activity (U/ml) for strains (□) LB-4 and (▨) LB-8 using 1% (w/v) forest products as carbon source. Error bars represent the mean \pm 2SD of duplicate samples.



Phylogenetic analysis

The isolate was identified on basis of 16S rRNA sequencing. A comparison of the DNA sequence with sequences in the National Center for Biotechnology Information (NCBI) database with BLAST software showed 99.99 % sequence identity with the published 16S rRNA sequences of *Bacillus pumilus*. The phylogenetic tree showed that strain LB-8 formed evolutionary lineage with members of the Bacillaceae family associated with members of the diverse *Bacillus* spectrum (Fig. 4). Several bacterial strains of *Bacillus*, *Pseudomonas*, *Clostridium* and *Cellulomonas* have shown cellulolytic activity under various culture conditions [22, 23]. Some strains belonging to this species have been reported for its ability of cellulase production, nevertheless, enzyme characterization reveals that this enzyme was optimally active at pH 6 and temperature 60 °C meanwhile the enzyme in this study is optimally active at pH 9 and temperature 50 °C [24].

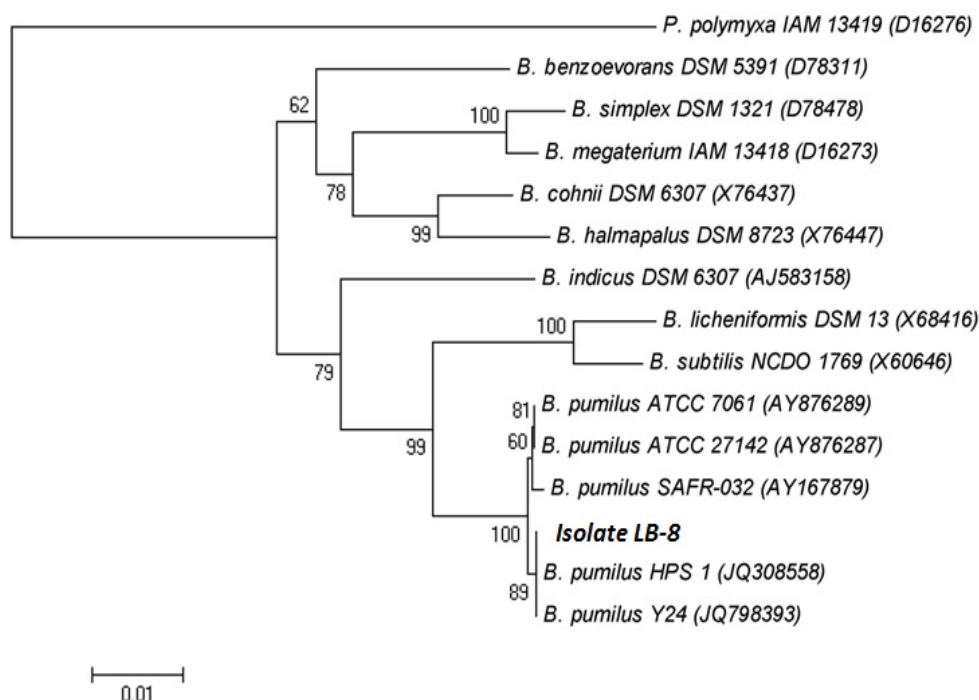


Figure43. Phylogenetic tree of members of the genus *Bacillus*, based on 16S rRNA gene sequences. The tree was constructed using the neighbor-joining method and the Jukes-Cantor distance matrix. *Paenibacillus polymyxa* IAM 13419 was used as the outgroup. Accession numbers are given in parentheses. Bar, genetic distance of 0.01.

CONCLUSION

Enzyme production can be improved in order to generate a suitable technology for economical processes. Different strategies have been implemented based in low-cost residues for cellulase production identifying major parameters affecting enzyme production [25-27]. The nine strains tested positive for cellulase production presented both halophilic and alkalotolerant properties. Strain LB-8 was able to growth at pH values ranging from 6 to 9; meanwhile positive cellular growth was observed at NaCl concentration up to 15 % (w/v). The use of agro-industrial waste like banana peel, sugarcane bagasse, rice husks, and wheat straw by strains LB-4 and LB-8 revealed better values of enzyme production from those values determined using forest products. This study presents several options of low-cost substrate to support cellulase production by these strains.

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REFERENCES

1. Gilkes, N.R., Kilburn, D.G., Miller, R.C., Warren, R.A.J. **1991**, Bacterial cellulases, *Bioresource Technology*, *36*(1), 21-35. DOI: 10.1016/0960-8524(91)90097-4
2. Amore, A., Pepe, O., Ventorino, V., Birolo, L., Giangrande, C., Faraco, V. **2013**, Industrial waste compost as source of novel cellulolytic strains and enzymes, *FEMS Microbiol. Lett.* *339*(2), 93-101. DOI: 10.1111/1574-6968.12057
3. Saini, J., Sini, R., Tewari, L. **2015**, Lignocellulosic agriculture waste as biomass feedstocks for second-generation bioethanol production: concepts and recent developments, *3 Biotech*, *5*, 337-353. DOI: 10.1007/s13205-014-0246-5
4. Gomes, D., Gama, M., Domingues, L. **2018**, Determinants on an efficient cellulase recycling process for the production of bioethanol from recycled paper sludge under high solid loadings, *Biotechnol. Biofuels*, *11*, 111. DOI: 10.1186/s13068-018-1103-2
5. Risacher, F. **1992**, Géochimie des lacs salés et croûtes de sel de l'Altiplano Bolivien, *Sci. Géol. Bull.* *45*, 135-214 (in French).
6. De Lourdes Moreno, M., Pérez, D., García, M.T., Mellado, E. **2013**, Halophilic bacteria as a source of novel hydrolytic enzymes, *Life*, *3*(1), 38-51. DOI: 10.3390/life3010038
7. Dos Santos, Q.Y., de Veras, O.B., de França, J.A.F., Gorlich-Lira, K., Velasques, J., Miggiolo, L., Dos Santos, A.E. **2018**, A new salt-tolerant thermostable cellulase from a marine *Bacillus* sp. strain, *J. Microbiol. Biotechnol.*, *28*(7), 1078-1085. DOI: 10.4014/jmb.1802.02037
8. Mann, H.B., Whitney, D.R. **1947** On a Test of Whether One of Two Random Variables Is Stochastically Larger than the Other. *An. Math. Stat.*, *18*, 50-60. DOI:10.1214/aoms/1177730491
9. Saitou, N., Nei, M. **1987**, The neighbor-joining method: a new method for reconstructing phylogenetic trees, *Mol. Biol. Evol.*, *4*(4), 406-425. DOI: 10.1093/oxfordjournals.molbev.a040454
10. Miller, G.M. **1959**, Use of dinitrosalicylic acid reagent for determination of reducing sugar, *Anal. Chem.*, *31*(3), 426-428. DOI: 10.1021/ac60147a030
11. Sangkharak, K., Vangsirikul, P., Janthachat, S. **2011**, Isolation of novel cellulase from agricultural soil and application for ethanol production, *Int. J. Adv. Biotechnol. Res.*, *2*(2), 230-239.
12. Ventosa, A., Nieto, J.J., Oren, A. **1998**, Biology of moderately halophilic aerobic bacteria, *Microbiol. Mol. Biol. Rev.*, *62*(2), 504-544. DOI: 10.1128/MMBR.62.2.504-544.1998
13. Horikoshi, K. **1996**, Alkaliphiles – from an industrial point of view, *FEMS Microbiol. Rev.*, *18*(2-3), 259-270.
14. Sahay, H., Mahfooz, S., Singh, A.K., Singh, S., Kaushik, R., Saxena, A.K., Arora, D.K. **2012**, Exploration and characterization of agriculturally and industrially important haloalkaliphilic bacteria from environmental samples of hypersaline Sambhar lake, India, *World J. Microb. Biotechnol.*, *28*, 3207-3217. DOI: 10.1007/s11274-012-1131-1
15. Ibrahim, A.S., Al-Salamah, A.A., Elbadawi, Y.B., El-Tayeb, M.A., Ibrahim, S.S.S. **2015**, Production of extracellular alkaline protease by new halotolerant alkaliphilic *Bacillus* sp. NPST-AK15 isolated from hyper saline soda lakes, *Electron J. Biotechnol.*, *18*(3), 236-243. DOI: 10.1016/j.ejbt.2015.04.001
16. Korany, A.H., Ali, A.E., Essam, T.M., Megahed, S.A. **2017**, Optimization of cellulase production by *Halobacillus* sp. QLS 31 isolated Lake Qarum, Egypt, *Appl. Biochem. Biotechnol.*, *183*, 189-199. DOI: 10.1007/s12010-017-2438-z
17. Cui, L., Meddeb-Mouelhi, F., Laframboise, F., Beauregard, M. **2015**, Effect of commercial cellulases and refining on kraft pulp properties: Correlation between treatment impacts and enzymatic activity components, *Carbohydr. Polym.*, *115*, 193-199. DOI: 10.1016/j.carbpol.2014.08.076
18. Kumar, N.V., Rani, M.E., Gunaseeli, R., Kannan, N.D. **2018**, Paper pulp modification and deinking efficiency of cellulase-xylanase complex from *Escherichia coli* SD5, *Int. J. Biol. Macromol.*, *111*, 289-295. DOI: 10.1016/j.ijbiomac.2017.12.126
19. Souza, L.T.A., Oliveira, J.S., Rodrigues, M.Q.R.B., Dos Santos, V.L., Pessela, B.C., Resende, R.R. **2015**, Macaúba (*Acroconia aculeata*) cake from biodiesel processing: a low-cost substrate to produce lipases from *Moniliella spathulata* R25L270 with potential application in the oleochemical industry, *Microb. Cell Fact.*, *14*(1), 87. DOI: 10.1186/s12934-015-0266-9
20. Awasthi, M.K., Wong, J.W.C., Kumar, S., Awasthi, S.K., Wang, Q., Wang, M., Ren, X., Zhao, J., Chen, H., Zhang, Z. **2018**, Biodegradation of food waste using microbial cultures producing thermostable α -amylase and cellulase under different pH and temperature, *Biores. Technol.*, *248*, 160-170. DOI: 10.1016/j.biortech.2017.06.160
21. Magalhaes, A., Carvalho, J.C., Melo Pereira, G.V., Karp, S. G. Camara, M.C., Medina, J.D.C., Soccol, C.R. **2019**, Lignocellulosic biomass from agro-industrial residues in South America: current developments and perspectives, *Biofuel Bioprod. Biorefin.*, *13*, 1505-1519. DOI: 10.1002/bbb.2048
22. Li, X., Yu, H.Y. **2012**, Purification and characterization of an organic-solvent-tolerant cellulase from a halotolerant isolate, *Bacillus* sp. L1, *J. In. Microbiol. Biotechnol.*, *39*(8), 1117-1124. DOI: 10.1007/s10295-012-1120-2
23. Kaushal, R., Sharma, N., Dogra, V. **2015**, Optimization of the production and molecular characterization of cellulase-free xylanase from an alkalophilic *Bacillus subtilis* SD8 isolated from paper mill effluent, *Appl. Biochem. Microbiol.*, *51*, 551-559. DOI: 10.1134/S0003683815050117
24. Ariffin, H., Abdullah, N., Umi, K.M.S., Shirai, Y., Hassan, M.A. **2006**, Production and characterization of cellulase by *Bacillus pumilus* EB3, *Int. J. Eng. Technol.*, *3*(1), 47-53.
25. Sakthivel, M., Karthikeyan, N., Jayaveny, R., Palani, P. **2010**, Optimization of culture conditions for the production of extracellular cellulase from *Corynebacterium lipophiloflavum*, *J. Ecobiotechnol.*, *2*(9), 6-13.
26. Zhao, C., Deng, Y., Wang, X., Li, Q., Huang, Y., Liu, B. **2014**, Identification and characterization of an anaerobic producing cellulolytic bacterial consortium from Great Basin hot springs with agricultural residues and energy crops, *J. Microbiol. Biotechnol.*, *24*(9), 1280-1290. DOI: 10.4014/jmb.1401.01022
27. Verma, N., Kumar, V., Bansal, M.C. **2020**, Comparative view on microbial consumption of agro-based lignocellulosic waste biomass in sustainable production of cellulases, *Biomass Conv. Bioref.* DOI: 10.1007/s13399-020-00617-0